

## SEQUENCE LISTING

- (1) GENERAL INFORMATION:
- (i) APPLICANT: Dolly, James Oliver
  Aoki, Kei Roger
  Wheeler, Larry Allen
  Garst, Michael Elwood
- (ii) TITLE OF INVENTION: MODIFICATION OF CLOSTRIDIAL TOXINS FOR USE AS TRANSPORT PROTEINS
- (iii) NUMBER OF SEQUENCES: 13
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Allergan, Inc.
  - (B) STREET: 2525 Dupont Drive
  - (C) CITY: Irvine
  - (D) STATE: CA
  - (E) COUNTRY: USA
  - (F) ZIP: 92623
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Diskette
  - (B) COMPUTER: IBM Compati/ble
  - (C) OPERATING SYSTEM: Windows 95
  - (D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION/DATA:
  - (A) APPLICATION NUMBER: 08/750,101
  - (B) FILING DATE: 01-MAY-1997
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: PCT/GB95/01253
  - (B) FILING DATE: 31-MAY-1995
  - (A) APPLICATION NUMBER: UK9410871.9
  - (B) FILING DATE / 31-MAY-1994
  - (A) APPLICATION NUMBER: UK9410871.1
  - (B) FILING DATE: 31-MAY-1994
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Fisher, Carlos A

July E

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| (B) REGISTRATION NUMBER: 36,510         |    |
|---|----|
| (C) REFERENCE/DOCKET NUMBER: 17044      |    |
| (ix) TELECOMMUNICATION INFORMATION:     |    |
| (A) TELEPHONE: 714-246-4920             |    |
| (B) TELEFAX: 714-246-4249               |    |
| (C) TELEX:                              |    |
|   |    |
| (2) INFORMATION FOR SEQ ID NO:1:        |    |
| (i) SEQUENCE CHARACTERISTICS:           |    |
| (A) LENGTH: 33 base pairs               |    |
| (B) TYPE: nucleic acid                  |    |
| (C) STRANDEDNESS: single                |    |
| (D) TOPOLOGY: linear                    |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: |    |
| GAGATGGTCG ACATGCCAAT AACCATAAAT AAT    | 33 |
| (2) INFORMATION FOR SEQ ID NO:2:        |    |
| (i) SEQUENCE CHARACTERISTICS:           |    |
| (A) LENGTH: 32 base pairs               |    |
| (B) TYPE: nucleic acid                  |    |
| (C) STRANDEDNESS: single                |    |
| (D) TOPOLOGY: linear                    |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: |    |
| ACGCGAAGCT TTTATCATGC AGTTCTATTA TA     | 32 |
| (2) INFORMATION FOR SEQ ID NO:3:        |    |
| (i) SEQUENCE CHARACTERISTICS:           |    |
| (A) LENGTH: 30 base pairs               |    |
| (B) TYPE: nucleic acid                  |    |
| (C) STRANDEDNESS: single                |    |
| (D) TOPOLOGY: linear                    |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: |    |
| FAGTACATGT ATAAGTGCGT GCATTAATAG        | 30 |
| (2) INFORMATION FOR SEQ ID NO:4:        |    |
| (i) SEQUENCE CHARACTERISTICS:           |    |
| (A) LENGTH: 20 base pairs               |    |

| <ul><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>   |    |
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| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:  |    |
| TTATACATGT ACTACATGGT  | 20 |
| (2) INFORMATION FOR SEQ ID NO:5:   |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 23 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul> |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:  |    |
| AAAGGCCTTT TGTTAATAAA CAA  | 23 |
| (2) INFORMATION FOR SEQ ID NO:6:   |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 26 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul> |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:  |    |
| GGAATTCTTA CTTATTGTAT CCTTTA 2   | 26 |
| (2) INFORMATION FOR SEQ ID NO:7:   |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul> |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:  |    |
| GCACATCAAC TTATACAT  | L8 |
| (2) INFORMATION FOR SEQ ID NO:8:   |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>   |    |

| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:   |    |
|---|----|
| ATGTATAAGT TGATGTGC   | 18 |
| (2) INFORMATION FOR SEQ ID NO:9:  |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>  |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:   |    |
| AACTTATATA TGCTGGAC   | 18 |
| (2) INFORMATION FOR SEQ ID NO:10:   |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>  |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:  |    |
| GTCCAGCATA TATAAGTT   | 18 |
| (2) INFORMATION FOR SEQ ID NO:11:   |    |
| <ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 0 amino acids</li> <li>(B) TYPE: amino acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> <li>(ii) MOLECULE TYPE: peptide</li> </ul> |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:  |    |
| Cys Ala Asn Gln Arg Ala Thr Lys Met Leu Gly Ser Gly 1 5 10  |    |
| (2) INFORMATION FOR SEQ ID NO:12:   |    |
| (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 29 base pairs  |    |

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

| <ul><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>   |    |
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| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:   |    |
| ATTTCACCAA TAACCATAAA TAATTTTAG  | 29 |
| (2) INFORMATION FOR SEQ ID NO:13:  |    |
| <ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 26 base pairs</li><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul> |    |
| (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:   |    |
| CGGGATCCTT CTGTATCATT GTAAAT   | 26 |
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